

1/10

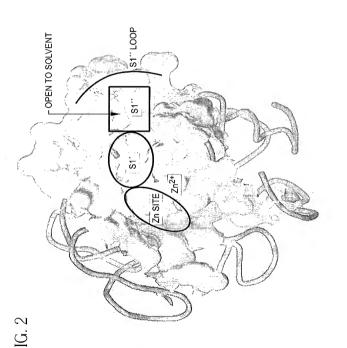
## FIG. 1

SEQUENCE LISTING <110> WARNER-LAMBERT <120> Matrix metalloproteinase inhibitors <130> A0000434 10 <140> <141> <160> 1 15 <170> PatentIn Ver. 2.1 <210> 1 <211> 471 <212> PRT 20 <213> Homo sapiens <400> 1 Met His Pro Gly Val Leu Ala Ala Phe Leu Phe Leu Ser Trp Thr His 25 Cys Arg Ala Leu Pro Leu Pro Ser Gly Gly Asp Glu Asp Asp Leu Ser Glu Glu Asp Leu Gln Phe Ala Glu Arg Tyr Leu Arg Ser Tyr Tyr His 30 Pro Thr Asn Leu Ala Gly Ile Leu Lys Glu Asn Ala Ala Ser Ser Met 35 Thr Glu Arg Leu Arg Glu Met Gln Ser Phe Phe Gly Leu Glu Val Thr Gly Lys Leu Asp Asp Asn Thr Leu Asp Val Met Lys Lys Pro Arg Cys 40 Gly Val Pro Asp Val Gly Glu Tyr Asn Val Phe Pro Arg Thr Leu Lys Trp Ser Lys Met Asn Leu Thr Tyr Arg Ile Val Asn Tyr Thr Pro Asp 115 120 125 45 Met Thr His Ser Glu Val Glu Lys Ala Phe Lys Lys Ala Phe Lys Val 135 Trp Ser Asp Val Thr Pro Leu Asn Phe Thr Arg Leu His Asp Gly Ile Ala Asp Ile Met Ile Ser Phe Gly Ile Lys Glu His Gly Asp Phe Tyr Pro Phe Asp Gly Pro Ser Gly Leu Leu Ala His Ala Phe Pro Pro Gly

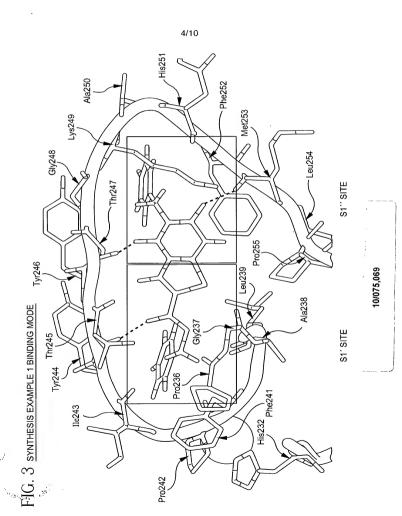
## 2/10

## FIG. 1A

_	Pr	o Ası	195	r Gl	y Gl	y As	p Al	a Hi:	s Ph	e Ası	o Ası	o As	205	u Th	r Tr	) Th
5	Ser	Ser 210	Ser	Lys	Gly	Tyr	Asn 215	Leu	Phe	Leu	Val	Ala 220	Ala	His	Glu	Phe
10	Gly 225	His	Ser	Leu	Gly	Leu 230	Asp	His	Ser	Lys	Asp 235	Pro	Gly	Ala	Leu	Met 240
	Phe	Pro	Ile	Tyr	Thr 245	Tyr	Thr	Gly	Lys	Ser 250	His	Phe	Met	Leu	Pro 255	Asp
15	Asp	Asp	Val	Gln 260	Gly	Ile	Gln	Ser	Leu 265	Tyr	Gly	Pro	Gly	Asp 270	Glu	Asp
20	Pro	Asn	Pro 275	Lys	His	Pro	Lys	Thr 280	Pro	Asp	Lys	Cys	Asp 285	Pro	Ser	Leu
	Ser	Leu 290	Asp	Ala	Ile	Thr	Ser 295	Leu	Arg	Gly	Glu	Thr 300	Met	Ile	Phe	Lys
25	Asp 305	Arg	Phe	Phe	Trp	Arg 310	Leu	His	Pro	Gln	Gln 315	Val	Asp	Ala	Glu	Leu 320
	Phe	Leu	Thr	Lys	Ser 325	Phe	Trp	Pro	Glu	Leu 330	Pro	Asn	Arg	Ile	Asp 335	Ala
30	Ala	Tyr	Glu	His 340	Pro	Ser	His	Asp	Leu 345	Ile	Phe	Ile	Phe	Arg 350	Gly	Arg
35	Lys	Phe	Trp 355	Ala	Leu	Asn	Gly	Tyr 360	Asp	Ile	Leu	Glu	Gly 365	Tyr	Pro	Lys
	Lys	Ile 370	Ser	Glu	Leu	Gly	Leu 375	Pro	Lys	Glu	Val	Lys 380	Lys	Ile	Ser	Ala
40	Ala 385	Val	His	Phe	Glu	Asp 390	Thr	Gly	Lys	Thr	Leu 395	Leu	Phe	Ser	Gly	Asn 400
	Gln	Val	Trp	Arg	Tyr 405	Asp	Asp	Thr	Asn	His 410	Ile	Met	Asp	Lys	Asp 415	Tyr
45	Pro	Arg	Leu	Ile 420	Glu	Glu	Asp	Phe	Pro 425	Gly	Ile	Gly	Asp	Lys 430	Val	Asp
50	Ala	Val	Tyr 435	Glu	Lys	Asn	Gly	Tyr 440	Ile	Tyr	Phe	Phe	Asn 445	Gly	Pro	Ile
		Phe 450	Glu	Tyr	Ser	Ile	Trp 455	Ser	Asn	Arg	Ile	Val 460	Arg	Val	Met	Pro
55.	Ala 465	Asn	Ser	Ile	Leu	Trp	Cys									



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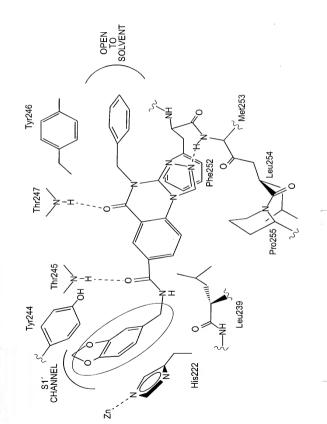


G. 4 SYNTHESIS EXAMPLE 1 BINDING MODE

 $FIG.\ \ 5$  synthesis example 10 binding mode

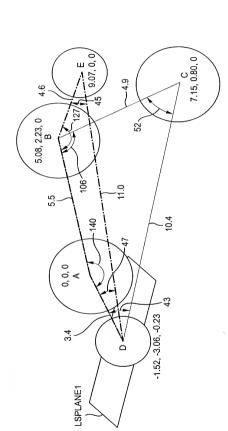
 ${\rm FIG.6}$ 

 $[G,\ 7]$  Synthesis example 39 binding mode



 $_{\rm T} FIG.~8~{\ }_{\rm SYNTHESIS}$  example 57 binding mode

10/10



9 COORDONATES IN THE SPACE OF THE HYDROPHOBIC GROUPS AND HYDROGEN BOND ACCEPTORS OF THE PHARMACOPHORE

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